

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/292,411

DATE: 11/28/2000  
TIME: 02:39:54

INPUT SET: S36148.raw

This Raw Listing contains the General  
Information Section and up to the first 5 pages.

SEQUENCE LISTING

(1) General Information:

ENTERED

(i) APPLICANT: Payne, David  
Lonsdale, John  
Milner, Peter  
Pearson, Stewart

(ii) TITLE OF INVENTION: FAB I

(iii) NUMBER OF SEQUENCES: 2

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Dechert Price & Rhoads  
(B) STREET: 4000 Bell Atlantic Tower, 1717 Arch Stre  
(C) CITY: Philadelphia  
(D) STATE: PA  
(E) COUNTRY: US  
(F) ZIP: 19103

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Diskette  
(B) COMPUTER: IBM Compatible  
(C) OPERATING SYSTEM: Windows 95  
(D) SOFTWARE: FastSEQ for Windows Version 2.0b

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 09/292,411  
(B) FILING DATE:  
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 08/790,043  
(B) FILING DATE:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Falk, Stephen T  
(B) REGISTRATION NUMBER: 36,795  
(C) REFERENCE/DOCKET NUMBER: GM50005

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 215-994-2488

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47 (B) TELEFAX: 215-994-2222  
48 (C) TELEX:  
49  
50

51 (2) INFORMATION FOR SEQ ID NO:1:  
52

53 (i) SEQUENCE CHARACTERISTICS:  
54 (A) LENGTH: 771 base pairs  
55 (B) TYPE: nucleic acid  
56 (C) STRANDEDNESS: double  
57 (D) TOPOLOGY: linear  
58  
59

60 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:  
61

62	ATGTTAAATC TTGAAAACAA AACATATGTC ATCATGGGAA TCGCTAATAA GCGTAGTATT	60
63	GCTTTTGGTG TCGCTAAAGT TTTAGATCAA TTAGGTGCTA AATTAGTATT TACTTACCGT	120
64	AAAGAACGTA GCCGTAAAGA GCTTGAAAAA TTATTAGAAC AATTAAATCA ACCAGAAGCG	180
65	CACTTATATC AAATTGATGT TCAAAGCGAT GAAGAGGTTA TTAATGGTTT TGAGCAAATT	240
66	GGTAAAGATG TTGGCAATAT TGATGGTGTA TATCATTCAA TCGCATTTGC TAATATGGAA	300
67	GACTTACGCG GACGCTTTTC TGAAACTTCA CGTGAAGGCT TCTTGTTAGC TCAAGACATT	360
68	AGTTCTTACT CATTACAAT TGTGGCTCAT GAAGCTAAAA AATTAATGCC AGAAGGTGGT	420
69	AGCATTGTTG CAACAACATA TTTAGGTGGC GAATTCGCAG TTCAAATTA TAATGTGATG	480
70	GGTGTGCTA AAGCGAGCTT AGAAGCAAAT GTTAAATATT TAGCATTAGA CTTAGGTCCT	540
71	GATAATATTC GCGTTAATGC AATTTAGCT GGTCCAATCC GTACATTAAG TGCAAAAGGT	600
72	GTGGGTGGTT TCAATACAAT TCTTAAAGAA ATCGAAGAGC GTGCACCTTT AAAACGTAAC	660
73	GTTGATCAAG TAGAAGTAGG TAAACAGCG GCTTACTTTR TAAGTGACTT ATCAAGTGCG	720
74	GTTACAGGTG AAAATATTCA TGTAGATAGC GGATTCCACG CAATTAAATA A	771

75  
76 (2) INFORMATION FOR SEQ ID NO:2:  
77

78 (i) SEQUENCE CHARACTERISTICS:  
79 (A) LENGTH: 256 amino acids  
80 (B) TYPE: amino acid  
81 (C) STRANDEDNESS: single  
82 (D) TOPOLOGY: linear  
83  
84

85 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:  
86

87	Met Leu Asn Leu Glu Asn Lys Thr Tyr Val Ile Met Gly Ile Ala Asn	
88	1 5 10 15	
89	Lys Arg Ser Ile Ala Phe Gly Val Ala Lys Val Leu Asp Gln Leu Gly	
90	20 25 30	
91	Ala Lys Leu Val Phe Thr Tyr Arg Lys Glu Arg Ser Arg Lys Glu Leu	
92	35 40 45	
93	Glu Lys Leu Leu Glu Gln Leu Asn Gln Pro Glu Ala His Leu Tyr Gln	
94	50 55 60	
95	Ile Asp Val Gln Ser Asp Glu Glu Val Ile Asn Gly Phe Glu Gln Ile	
96	65 70 75 80	
97	Gly Lys Asp Val Gly Asn Ile Asp Gly Val Tyr His Ser Ile Ala Phe	
98	85 90 95	
99	Ala Asn Met Glu Asp Leu Arg Gly Arg Phe Ser Glu Thr Ser Arg Glu	

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION** *US/09/292,411*

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**INPUT SET: S36148.raw**[illegible]

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**SEQUENCE VERIFICATION REPORT**  
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Line

Error

Original Text

PAGE: 1

**SEQUENCE MISSING ITEM REPORT**  
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< < THERE ARE NO ITEMS MISSING > >

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**SEQUENCE CORRECTION REPORT**  
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